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Author contact	<p>marlies.coopman@kuleuven-kulak.be</p> <p>+32 56 24 62 48</p>
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Context dependency of infectious disease: the cyanobacterium *Microcystis aeruginosa* decreases White Bacterial Disease in *Daphnia magna*.

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1 **Context dependency of infectious disease: the cyanobacterium *Microcystis aeruginosa***
2 **decreases White Bacterial Disease in *Daphnia magna*.**

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4 MARLIES COOPMAN, KOENRAAD MUYLEAERT, BENJAMIN LANGE, LIEN REYSERHOVE AND
5 ELLEN DECAESTECKER

6
7 *Aquatic Biology, KULeuven-Kulak, Kortrijk, Belgium*

8
9 Correspondence: Marlies Coopman, Aquatic Biology, KULeuven-Kulak, 53 Etienne Sabbelaan,
10 8500 Kortrijk, Belgium. E-mail: marlies.coopman@kuleuven-kulak.be

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12 *Running headline: Microcystis decreases parasitism in Daphnia*

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14 *Keywords: cyanobacteria, Daphnia, parasitism, antibacterial effect, toxic metabolites*

Summary

1. We investigate whether an increase in the occurrence of cyanobacterial blooms affects zooplankton-parasite interactions. Cyanobacteria are expected to be of poor food quality for zooplankton hosts and are therefore expected to increase parasitism. Nevertheless, simultaneous exposure to both stressors may lead to different results, given the antibacterial secondary metabolites of cyanobacteria.
2. We exposed the zooplankter *Daphnia magna* to the cyanobacterial species *Microcystis aeruginosa* and the parasite that causes White Bacterial Disease in *D. magna*. Increased *M. aeruginosa* concentrations reduced the percentage of infected individuals and as such protected *D. magna* against parasitism. Interactions between *M. aeruginosa* and the parasite were antagonistic in terms of percentage of surviving *Daphnia*, total offspring per female and clutch size. Additional plating experiments showed a direct negative effect of *Microcystis* on bacterial growth.
3. The results suggest that changes in phytoplankton affect host-parasite interactions in zooplankton. Contrary to the prevailing paradigm that multiple stressors often induce additive or synergistic effects, we report an antagonistic effect of the presence of cyanobacterial stress on parasites in *Daphnia*. Thus, assessment of the outcome of host-parasite interactions needs to incorporate the environmental context.

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34 **Introduction**

35 The intensification of human activities has profoundly affected biotic interactions in
36 natural ecosystems. Often overlooked, host-parasite interactions are important in this
37 context. In the long term, these interactions fuel biodiversity (Mouritsen & Poulin, 2005).
38 and we need to understand how environmental changes affect host-parasite dynamics
39 (Johnson & Carpenter; 2008, Lennon & Martiny, 2008; Duffy *et al.*, 2012). With respect to
40 ecological effects on immunity, enhanced stressful conditions have often been associated
41 with immunosuppression (Acevedo-Whitehouse & Duffus, 2009; Schmid-Hempel, 2011;
42 Adamo, 2012). On the other hand, environmental changes that do not involve
43 immunological responses may counteract the need for immune-based host responses
44 (Parker *et al.*, 2011). There is thus a need to address the context dependency of infectious
45 diseases (Ibelings *et al.*, 2011; Rohr *et al.*, 2011; Overholt *et al.*, 2012).

46 In this paper, we aim to unravel the effect of the cyanobacterium *Microcystis*
47 *aeruginosa* on *Daphnia magna* parasitism. Cyanobacterial blooms have become a worldwide
48 problem in many freshwater ecosystems (Codd, Morrison & Metcalf, 2005; Paerl & Huisman,
49 2008; Davis *et al.*, 2009; Kosten *et al.*, 2012). The negative effect of cyanobacteria on
50 zooplankton and especially on *Daphnia* species is well documented (Ferro, Azevedo &
51 DeMott, 2000; Asselman *et al.*, 2012; Lemaire *et al.*, 2012). Many previous studies have
52 focused on detrimental, toxic effects of microcystins (Demott, Zhang & Carmichael, 1991;
53 Reinikainen, Ketola & Walls, 1994; Rohrlack *et al.*, 1999). However, not all negative effects
54 are attributed to the presence of toxins (Berry *et al.*, 2008); recent studies also attribute it to
55 the effects of low food quality (Martin-Creuzburg, von Elert & Hoffmann, 2008; von Elert, Zitt
56 & Schwarzenberger, 2012). Cyanobacteria are low quality food for zooplankton in terms of
57 biochemical composition because of the absence of long chain poly-unsaturated fatty acids

(PUFAs) and sterols (Muller-Navarra *et al.*, 2000; von Elert, Martin-Creuzburg & Le Coz, 2003; Martin-Creuzburg & von Elert, 2009) and the presence of protease inhibitors (Namikoshi & Rinehart, 1996; Rohrlack *et al.*, 2004; Schwarzenberger *et al.*, 2010).

Our focus here is the effect of food quality on zooplankton-parasite interactions and we therefore use a non-microcystin producing strain of *M. aeruginosa*. As well as their low food quality, cyanobacteria also harbour diverse secondary metabolites, including lipopeptides, amino acids, fatty acids, macrolides, amides that exhibit a wide range of bioactivities (e.g. antibacterial, antifungal, antialgal, antiprotozoan, antiviral) (Abed, Dobretsov & Sudesh, 2009; Singh *et al.*, 2011). Such secondary metabolites with biological activity have been found in *M. aeruginosa* (Ishida *et al.*, 1997; Abed *et al.*, 2009; Pradhan *et al.*, 2011). Nevertheless, *in vivo* experiments studying the effects on parasitism within hosts are extremely rare.

Our study animal is *Daphnia*, a keystone player in lake ecosystems because of its grazing effects on phytoplankton; *Daphnia* is also a laboratory model system (Miner *et al.*, 2012). Specifically, our study involves *Daphnia magna* because of its strong responses to changes in nutrient enrichment (Verreydt *et al.*, 2012) and associated *Microcystis* blooms. The responses of *Daphnia magna* to *Microcystis* exposure are well described (Nizan, Dimentman & Shilo, 1986; Schwarzenberger *et al.*, 2010; Lemaire *et al.*, 2012). *Daphnia magna* is also a model species for host-parasite interactions (Ebert, 2005; Ebert, 2011). The parasite we used causes White Bacterial Disease (WBD) and was selected for its strong virulence effects and its common occurrence in *Daphnia* populations in Western and Northern Europe (Decaestecker *et al.*, 2005; Ebert, 2005). Relatively few studies have investigated the role of food quality on *Daphnia* parasitism, but increasing evidence shows that changes in essential nutrients in food affect *Daphnia*-parasite interactions (Frost, Ebert

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82 & Smith, 2008; Hall *et al.*, 2009; Frost *et al.*, 2010; Stjernman & Little, 2011; Aalto &
83 Pulkkinen, 2013). *Daphnia* that are limited by food availability are considered to be more
84 susceptible to parasites (Frost *et al.*, 2008; Schoebel, Wolinska & Spaak, 2010; Vale *et al.*,
85 2011). Given that cyanobacteria are considered to be of poor food quality (Muller-Navarra *et*
86 *al.*, 2000; von Elert *et al.*, 2003; Martin-Creuzburg & von Elert, 2009), susceptibility of
87 *Daphnia* to its parasite upon cyanobacterial exposure is expected to increase. The prevailing
88 paradigm in multiple-stressor ecology predicts an increase in disease infectivity or virulence
89 via weakened, immunosuppressed hosts (as detected in Coors *et al.*, 2008; Jansen *et al.*,
90 2011). Nevertheless, given the antibacterial effects of secondary metabolites detected *in*
91 *vitro* in cyanobacteria, a decrease in bacterial parasitism would not be an unlikely outcome.

92

93 **Methods**

94 **Isolation of *D. magna*, parasites and *M. aeruginosa***

95 We investigated the impact of a non-microcystin producing *M. aeruginosa* strain on
96 *D. magna* upon exposure to the parasite that causes WBD. Due to increasing interest in non-
97 microcystin related characteristics of cyanobacteria, such as food quality and the presence of
98 bioactive secondary metabolites, a single celled, non-microcystin producing cyanobacterial
99 strain of *M. aeruginosa* was used to unravel the cyanobacterial effect in the absence of toxic
100 microcystins (*M. aeruginosa* CCAP1450/1, provided by the Culture Collection of Algae and
101 Protozoa, UK, and isolated in 1948 from the highly productive Lake Mendota, Wisconsin,
102 USA, where *M. aeruginosa* regularly dominates the lake) (Soranno, 1997). The strain was
103 confirmed not to contain any microcystins. We performed polymerase chain reaction (PCR)
104 amplification of three crucial genes of the microcystin synthetase cluster (mcyA, mcyB and
105 mcyE: Neilan *et al.*, 1999; Hisbergues *et al.*, 2003; Vaitomaa *et al.*, 2003), but no amplified
106 products were detected. The microcystin-producing *Microcystis aeruginosa* PCC7806 was
107 used as a positive control. This strain is also available in other culture collections under
108 different names (*M. aeruginosa* 1036, 1036AX, NIVA-CYA43, SAG 1450-1, UTCC 73, UTEX
109 LB2061, UTEX 2061 *Microcystis* sp. PCC 7005 and *Synechocystis* sp. PCC 7005 and ATCC
110 27153) and different research groups have demonstrated its inability to produce
111 microcystins (reviewed by Lyra *et al.*, 2001). *M. aeruginosa* strains were cultured in modified
112 WC medium in 1 L bottles that were aerated and stirred (Guillard & Lorenzen, 1972, the
113 medium did not contain Tris). Stocks of *D. magna*, *M. aeruginosa* and *Scenedesmus obliquus*
114 were cultured in a climate chamber at 20±2°C with a light:dark cycle of 16:8 h. Experiments
115 were performed under the same conditions.

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116 The *D. magna* clone, originally obtained by hatching dormant eggs, was isolated from
117 the sediment of a pond in Heverlee, Belgium (OM2, “Abdij van ‘t Park”, 50°51'47.67"N;
118 4°43'16.36"E). This *D. magna* population has been well described with respect to its parasitic
119 interactions (Decaestecker *et al.*, 2005; Decaestecker *et al.*, 2007; Decaestecker *et al.*, 2013).
120 *Daphnia* clone 16.1 was chosen because of its good performance in laboratory conditions
121 and its sensitivity to the parasite. Before running the experiment, maternal effects were
122 controlled by culturing fifteen iso-female lines separately under standardized conditions
123 (20±2°C with a light:dark cycle of 16:8h) for three generations. Lineages were fed daily with
124 1.5 mg C L⁻¹ *Scenedesmus obliquus* and medium was refreshed three times a week. *Daphnia*
125 culturing and experiments were performed in ADaM medium (Kluttgen *et al.*, 1994).

126 We used the parasite that causes White Bacterial Disease (WBD) in *D. magna*. WBD,
127 also known as White Fat Cell Disease (WFD), is caused by a small coccoid parasite, most
128 likely a bacterium that infects the adipose tissue, resulting in a bright white reflection with a
129 greenish shine from infected tissue that is only visible in reflected light (Ebert, 2005). The
130 infection is virulent and is transmitted horizontally (Decaestecker *et al.*, 2005). Infected
131 females were collected from another *D. magna* population than the clone that was used in
132 the final experiment, to exclude pre-adaptation. The parasite was isolated from a pond in
133 Zonhoven, Belgium (lake 21, 50° 59' 12.72" N en 5° 20' 35.56" E) and kept in a 400 L
134 container with non-infected *D. magna* individuals from the same pond. Upon establishment
135 of infections, infected *Daphnia* individuals were picked from the 400 L container and kept at
136 4°C until the experiment was performed. In the parasite treatment, *Daphnia* were exposed
137 to a solution of homogenized infected *Daphnia* (3x25 cadavers in 3x5 mL). Parasite exposure
138 consisted of one infected individual for three females to be infected. An equal number of
139 healthy individuals was homogenized for use in the control treatment.

140 ***In vivo Daphnia* exposure to *Microcystis* and WBD**

141 Neonates (<24 h old) from the second brood of the third generation of the maternal
142 lines were isolated and five individuals were pooled and placed in a 200 mL experimental jar
143 (design according to Lemaire *et al.*, 2012). Five replicates were performed per treatment
144 combination, each from a different mother line to control for maternal effects and
145 interdependency of the different replicates. Neonates were fed daily 5 mg C L⁻¹ of a mixture
146 of *M. aeruginosa* and *S. obliquus* in concentrations of 0%, 10%, 20%, 50%, 70% and 100% *M.*
147 *aeruginosa*. This scale was chosen based on results of earlier pilot experiments. In total, 60
148 experimental jars were involved (6 concentrations x 2 parasite treatments (= with and
149 without parasite) x 5 replicates). The medium was refreshed daily until day 4. From day five,
150 female *D. magna* juveniles were exposed for five days to a mixture of homogenized WBD-
151 infected *Daphnia* tissue. On day five, the *Daphnia* were exposed to the parasite or placebo
152 solution in 50 mL, at day six and seven in 100 mL and at day eight and nine in 200 mL.
153 Addition of the parasite or placebo solution was spread over three consecutive days and the
154 added volume was recalculated every day to make sure that each living *Daphnia* was
155 exposed to an equal number of parasites. From day ten until the end of the experiment, the
156 medium was refreshed daily. Survival was measured daily as the number of surviving
157 individuals out of five *Daphnia* in each jar. The number of offspring produced and the
158 number of WBD-infected individuals were recorded daily. On day 21, two females from each
159 microcosm were randomly chosen, measured (as in Coors & De Meester, 2008) and their
160 clutch size was recorded.

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In vitro* test of the antibacterial effect of *Microcystis

To confirm a potential direct negative effect of the cyanobacterial food on the *D. magna* parasite, we tested antimicrobial activity of the extracts of *M. aeruginosa* and *S. obliquus* using a paper disc diffusion assay. As no standard cultures and plating techniques are yet available for WBD, plating experiments were carried out using *Escherichia coli* (NovaBlue Singles™ Competent Cells, Novagen, EMD Biosciences, Inc, Darmstadt, Germany). 10 mg C of the *M. aeruginosa* or *S. obliquus* strain was concentrated by centrifugation and lysed using five freeze-thaw cycles. Sterilized filter paper discs (6 mm) were saturated with the filtered extract (0.22 µm) and placed on a monolayer of *E. coli* on agar. Paper discs saturated with a 3 mg mL⁻¹ tetracycline solution were used as positive controls. Cultures were incubated for 20 hours at 37°C before diameters of the inhibition zones surrounding the disc were measured (Andrews, 2001).

Statistical analysis

Generalized Linear Models (GLMs) were used to investigate the effect of the parasite exposure and the effect of increasing *M. aeruginosa* concentration (0%, 10%, 20%, 50%, 70%, 100%, categorical data). Percentage surviving *Daphnia* was analyzed as binomially-distributed data, as the ratio of surviving *Daphnia* to the number of dead *Daphnia*. Body size data (mean value for each experimental jar) were analyzed as normally-distributed data. Total offspring per female was calculated as the sum of daily offspring per jar divided by total surviving *Daphnia*, since all *Daphnia* in this experiment were female. Total offspring per female was also analyzed as normally-distributed data. Clutch size (eggs per female at the end of the experiment) was treated as count data and analyzed by a Poisson distribution (mean value for each experimental jar). Time to first clutch (i.e. the time when the first

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3 186 juveniles appeared in the jar) was analyzed by survival analysis, specifically the Cox
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5 187 proportional hazards model (Efron method). In this analysis, surviving females that had not
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7 188 reproduced by the end of the experiment were taken into account as right-censored data.
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10 189 Percentage infected *Daphnia* per jar was analyzed as binomially-distributed data. In all
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12 190 analyses the ratio of the residual deviance to the degrees of freedom was verified to be one.
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14 191 If not, data were corrected for over- or underdispersion. Then the dispersion parameter was
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16 192 not fixed at one, but estimated by moments. To compare the influence of the parasite
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18 193 (control versus exposed) at different *S. obliquus*/*M. aeruginosa* mixtures, a Tukey post hoc
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20 194 analysis was performed for the response variables percentage surviving *Daphnia*, body size,
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22 195 clutch size, time to first clutch and total offspring per female.
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26 196 No statistical analyses were performed on the antibacterial activity of tetracycline
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28 197 and the lysates of *Microcystis* and *Scenedesmus* since only the appearance of a halo,
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30 198 showing antibacterial effects of the solution on *E. coli*, was important for this experiment. All
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32 199 tests were performed using the free statistical tool pack R version 2.10. ([http://www.r-](http://www.r-project.org/)
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34 200 [project.org/](http://www.r-project.org/)).
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Results

The results of our experiment almost always showed a significant negative effect when the two stressors were applied separately. *M. aeruginosa* negatively affected the *D. magna* clone: the percentage of surviving *Daphnia*, body size and clutch size decreased (Fig. 1a,b,c, Table 1) while time to first clutch increased with an increasing *M. aeruginosa* concentration (Fig. 1d, Table 1), resulting in a significant decrease in total offspring per female with increasing *M. aeruginosa* concentration (Fig 1e, Table 1). Exposure to WBD had a negative effect on *Daphnia* by reducing the percentage of surviving *Daphnia* (marginally significant) and clutch size (Fig. 1a,c, Table 1) but had a positive effect on *Daphnia* body size and total offspring per female (Fig. 1b,e Tables 1). No parasite effect was observed for time to first clutch (Fig. 1d, Table 1).

When *D. magna* was simultaneously exposed to both stressors, interaction effects (i.e. antagonistic effects between the two stressors) could be detected, especially between 0% and 20% *M. aeruginosa*. There were significant interaction effects between the *M. aeruginosa* and WBD treatment on *Daphnia* survival (percentage surviving *Daphnia*), total offspring per female and clutch size (Table 1). There was no interaction effect for body size or time to first clutch (Tables 1). The negative effect of WBD on percentage *Daphnia* survival and clutch size was present when no *M. aeruginosa* were present in the food treatment, but disappeared at 20% *M. aeruginosa* (Fig. 1a,c, Table 2).

No interaction was found for time to first clutch over the range of 0 to 100% *M. aeruginosa* but a reduction of the negative *Microcystis* effect was observed upon simultaneous WBD exposure and 20% *M. aeruginosa* in the food treatment (Fig. 1d, Table 2). Non-WBD exposed *Daphnia* produced their first brood later when fed with 20% *M. aeruginosa* in comparison with the 0% *M. aeruginosa* food treatment (Tukey post hoc

analysis of non-WBD exposed *Daphnia* between 0% and 20% *Microcystis*: $p = 0.025$). Upon exposure to WBD, there was no difference in time to first brood between the 0% and the 20% *M. aeruginosa* food treatments (Tukey post hoc analysis of WBD exposed *Daphnia* between 0% and 20% *Microcystis*: $p = 0.984$). Further, total offspring per female was higher in the WBD exposed assay compared to the placebo treatment at 20% *M. aeruginosa* in the food treatment (Fig. 1e, Table 2). This is probably due to the earlier first brood at 20% *M. aeruginosa* (Fig. 1d, Table 2).

There was a reduction in the percentage infected *Daphnia* upon increasing *M. aeruginosa* concentration in the food (Fig. 1f, GLM for binomially-distributed data: likelihood ratio 24.464, degrees of freedom 5, $p < 0.001$). Additionally, a paper disc diffusion bioassay demonstrated direct antibacterial activity of the *M. aeruginosa* lysate on *E. coli* (Table 3). The lysate of *S. obliquus* did not show this activity.

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Discussion

Both the cyanobacterial strain *M. aeruginosa* and the parasite WBD individually induced negative fitness effects in *D. magna* and can thus be considered as biotic stressors. In view of its negative effects on life history traits, *M. aeruginosa* has generally been considered to be a biotic stressor for *D. magna* (Ferro et al., 2000; Asselman et al., 2012; Lemaire et al., 2012). The present study confirms that these negative effects are indeed present upon individual exposure of *Daphnia* to *M. aeruginosa*. The negative effects associated with *M. aeruginosa* can be assigned to toxic substances such as microcystins (Demott et al., 1991) or gamma-linolenic acid (Reinikainen et al., 2001) or poor food quality such as the absence of sterols and certain essential PUFAs (Martin-Creuzburg et al., 2008) or the presence of protease inhibitors (von Elert et al., 2012). The frequently observed mechanical obstruction of *Daphnia*'s filtering apparatus (Lampert, 1987; De Bernardi & Giussani, 1990; DeMott, Gulati & Van Donk, 2001) does not apply here because the *Microcystis* strain that we used consisted only of single cells.

We detected an antagonistic combined effect of the stress of *M. aeruginosa* and WBD in this particular association of host clone and parasite strain. This is contrary to our original expectation based on the prevailing paradigm that multiple stressors (especially when toxicants are involved) are assumed to induce mostly additive or synergistic effects (Coors et al., 2008; Sures, 2008; Holmstrup et al., 2010). *Microcystis* and cyanobacterial stress in general can be considered as an exception, given the antibacterial properties of secondary metabolites, which have so far mainly been investigated *in vitro* (Ishida et al., 1997; Abed et al., 2009; Pradhan et al., 2011). One research group has detected negative effects of cyanobacteria on parasitism in tomato plants *in vivo* (Khan et al., 2005; Khan et al., 2007). Antibacterial effects may make cyanobacteria an attractive source for

pharmacologically active compounds (reviewed by Abed *et al.*, 2009; Rastogi & Sinha, 2009). The production of such substances may confer a survival advantage by aiding the cyanobacterium in competition with other bacteria or algae (Ishida *et al.*, 1997; Singh *et al.*, 2001; Leflaive & Ten-Hage, 2007). Note that our *M. aeruginosa* strain has been cultured for almost seventy years in a laboratory. In this strain, competition with other algae or cyanobacteria is absent, but other bacteria may be present (non-axenic culture, see Culture Collection of Algae and Protozoa, UK). However, as only the lysate was tested *in vitro* against *E. coli*, the specific substances that are responsible for the antibacterial effect are not known.

Our study also revealed a negative effect of *Microcystis* on the parasite. This antibacterial result cannot, however, be generalized because our *in vivo* study is based on a single parasite species-*Microcystis* strain exposed to a single *D. magna* clone. The *in vitro* response did, however, also show an effect of the *M. aeruginosa* strain on *E. coli*. We therefore hypothesize that feeding on antibacterial cyanobacteria may help reduce infections in *Daphnia* in the field. The parasite that causes White Bacterial Disease has been found in ponds where *M. aeruginosa* blooms occur (OM2, Heverlee, Belgium), but to our knowledge co-occurrence of *M. aeruginosa* and WBD has never been reported.

Our experiment has also demonstrated that, upon exposure to both stressors, *Daphnia* begins reproducing earlier. While fecundity compensation through a reproduction shift upon infection had been detected in *Daphnia* before (Ebert *et al.*, 2004; Chadwick & Little, 2005), the fecundity compensation reported here only occurred when both stressors were present. The time to start reproducing was not delayed in the presence of both stressors as was the case upon *M. aeruginosa* exposure only. Together with an increase in clutch size, which was probably due to a decreasing parasite effect, this resulted in an

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287 increase in total offspring per surviving female.

288 Rather than being an enhanced immune response elicited in *Daphnia* by the presence

289 of cyanobacteria, the detected antibacterial activity is more likely a direct effect of the

290 cyanobacteria on the WBD parasite. Nevertheless, we cannot exclude a role for enhanced

291 immuno-competence in *Daphnia* (immuno-enhancement has been reported upon short

292 stress exposure; Demas, Adamo & French, 2011). Much attention has been devoted in the

293 past decade to the effects of environmental constraints on the role immune systems play in

294 determining host fitness in the wild (Lochmiller & Deerenberg, 2000; Woodhams *et al.*, 2008;

295 Sadd & Schmid-Hempel, 2009). In general, and with respect to invertebrates, immuno-

296 suppression is expected in stressful conditions because of reduced investment in mounting

297 an immune response (Schmid-Hempel, 2011; Adamo, 2012). In this context it is interesting

298 to speculate whether the gut microbiota of *Daphnia* may also have mediated the

299 antibacterial response (Rengpipat *et al.*, 2000; Dillon *et al.*, 2005) or have stimulated

300 mortality in infected *Daphnia*.

301 It is known that *Daphnia* can avoid *Microcystis*, probably based on taste (Rohrback *et*

302 *al.*, 2001; Tillmanns, Burton & Pick, 2011). In the presence of single and non-microcystin

303 producing *Microcystis* cells, *Daphnia* reduces its ingestion rate (Tillmanns *et al.*, 2011). In

304 addition to low food quality of *Microcystis*, this may help explain reduced survival in the

305 absence of the parasites. Consequently, as spores of WBD are most likely taken up by the

306 filter apparatus of *Daphnia*, the ingestion of WBD spores will also be reduced in the presence

307 of *Microcystis*. This may explain the decline in the percentage infected *Daphnia* upon

308 increasing *M. aeruginosa* concentration in the food. However, other factors may also come

309 into play, as the reduced ingestion rate may not completely explain *Daphnia* survival when

310 exposed to WBD. At 20% *Microcystis*, food and spores can still be filtered and *Daphnia* still

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3 311 become infected, but the consequent death penalty disappears from this concentration
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5 312 onwards. It seems that infection is kept under control from 20% *Microcystis* onwards.
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7 313 *Daphnia* still get infected, but survive the infection. It is possible that *Daphnia* develops
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9 314 resistance against *Microcystis* and is therefore also more defended against parasite spores
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11 315 (cross-reactivity, Schmid-Hempel, 2011). Another explanation could be the suppression of
12
13 316 disease by the antibacterial metabolites produced by *Microcystis* (and an associated
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15 317 reduction in virulence through a reduction in bacterial growth). However, as decreased
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17 318 infection rates were also found with phosphorus-limited food (Frost *et al.*, 2008) follow-up
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19 319 studies are needed to disentangle the effects of low food quality, immune host responses
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21 320 and the direct effect of antibacterial substances.
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554 **Tables**

555 **Table 1** Results of Generalized Linear Models (GLM) testing for the effect of parasite
 556 exposure (P) and *M. aeruginosa* concentration (C) in the food on percentage surviving
 557 *Daphnia*, body size, clutch size and total offspring per female. Time to first clutch was
 558 analyzed by a Cox proportional hazards model.

559

Response variable	Effect	Likelihood ratio	Degrees of Freedom	p-value
Percentage surviving <i>Daphnia</i>	C	49.376	5	<0.001
	P	13.647	1	<0.001
	P x C	18.616	5	0.002
Body size	C	55.366	5	<0.001
	P	7.760	1	<0.001
	P x C	2.807	5	0.730
Clutch size	C	33.184	5	<0.001
	P	43.319	1	<0.001
	P x C	23.611	5	<0.001
Time to first clutch	C	33.065	5	<0.001
	P	1.135	1	0.287
	P x C	9.048	5	0.107
Total offspring per female	C	119.060	5	<0.001
	P	7.592	1	0.006
	P x C	24.297	5	<0.001

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Table 2 Results of Tukey post hoc tests analyzing different response variables between WBD exposed and non-exposed *D. magna* females fed with mixtures of 0/100% and 20/80% *M. aeruginosa*/*S. obliquus*.

Response variable	Percentage <i>Microcystis</i>	Estimated value	Standard error	Z-value	p-value
Percentage surviving	0%	3.932	1.516	2.593	0.056
<i>Daphnia</i>	20%	0.399	0.870	0.459	0.998
Body size	0%	0.378	0.136	2.786	0.032
	20%	0.355	0.128	2.775	0.033
Clutch size	0%	-2.152	0.433	-4.973	<0.001
	20%	-0.331	0.215	-1.540	0.547
Time to first clutch	0%	-0.748	0.703	-1.064	0.867
	20%	2.452	0.756	3.243	0.007
Total offspring per	0%	-8.900	3.230	-2.755	0.035
female	20%	11.413	3.230	3.533	0.002

Table 3 Antibacterial activity of tetracycline (3 mg mL⁻¹) and the lysate of *Scenedesmus* and *Microcystis* against *E. coli* using the disc diffusion bioassay.

Substrate	Inhibition zone
Tetracycline	YES
Lysate of <i>S. obliquus</i>	NO
Lysate of <i>M. aeruginosa</i>	YES

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573 **Figure legends**

574 **Fig. 1** (a) Percentage surviving *Daphnia*, (b) body size and (c) clutch size of *Daphnia* surviving

575 on day 21 of the experiment, (d) time to first clutch, (e) total offspring per female, and (f)

576 percentage of *D. magna* females infected with White Bacterial Disease as a function of the

577 *M. aeruginosa* - *S. obliquus* mixtures. Means are given and error bars indicate one standard

578 error.

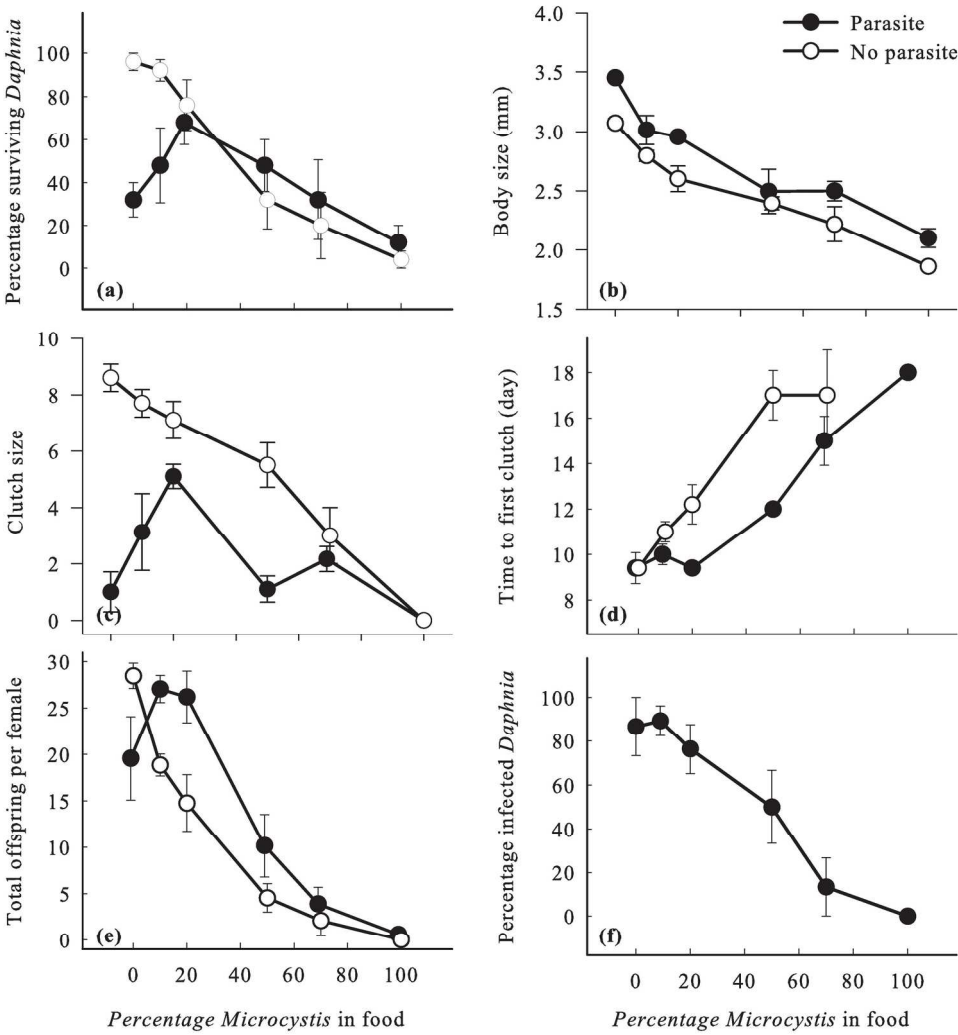
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580 **Illustrations**

581 Fig. 1 See attachment

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